SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Hartley, James L. Brasch, Michael A.
- (ii) TITLE OF INVENTION: Recombinational Cloning Using Engineered Recombination Sites
- (iii) NUMBER OF SEQUENCES: 35
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX, P.L.L.C.
 - (B) STREET: 1100 New York Ave., N. W. Suite 600
 - (C) CITY: Washington (D) STATE: DC
 - (E) COUNTRY: USA
 - (F) ZIP: 20005-3934
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: (To be assigned)
 - (B) FILING DATE: (Herewith)
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 09/233,493
 - (B) FILING DATE: 20-JAN-1999 (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 09/005,476
 - (B) FILING DATE: 12-JAN-1998
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/663,002
 - (B) FILING DATE: 07-JUN-1996
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/486,139
 - (B) FILING DATE: 07-JUN-1995
 - (C) CLASSIFICATION:
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 202-371-2600 (B) TELEFAX: 202-371-2540
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25 base pairs
 - (B) TYPE: nucleic acid

(C) STRANDEDNESS: both (D) TOPOLOGY: both

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

(ii) MOLECULE TYPE: cDNA

(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both	
(A) LENGTH: 25 base pairs (B) TYPE: nucleic acid	
(D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(x±) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
AGCCWGCTTT YKTRTACNAA CTSGB 2	5
(2) INFORMATION FOR SEQ ID NO:3:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GTTCAGCTTT CKTRTACNAA CTSGB 2.	5
(2) INFORMATION FOR SEQ ID NO:4:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii' MOLECULE TYPE: cDNA	
(x1) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
AGCCWGCTTT CKTRTACNAA GTSGB 2:	5

	5	
(2)	INFORMATION FOR SEQ ID NO:5:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
GTT	CAGCTTT YKTRTACNAA GTSGB	25
(2)	INFORMATION FOR SEQ ID NO:6:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
	(ii) MOLECULE TYPE: eDNA	
	(x^{\pm}) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
7000	CTECTIT TITGTACAAA CITGI	
		25
(2)	INFORMATION FOR SEQ ID NO:7:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
AGC	CTGCTTT CTTGTACAAA CTTGT	25
(2)	INFORMATION FOR SEQ ID NO:8:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
	(ii) MOLECULE TYPE: cDNA	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
ACCCAGCTTT CTTGTACAAA CTTGT	25
(2) INFORMATION FOR SEQ ID NO:9:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
GTTCAGCTTT TTTGTACAAA CTTGT	25
(2) INFORMATION FOR SEQ ID NO:10:	-
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
GTTCAGCTTT CTTGTACAAA CTTGT	25
(2) INFORMATION FOR SEQ ID NO:11:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
GTTCAGCTTT CTTGTACAAA GTTGG	2.5
(2) INFORMATION FOR SEQ ID NO:12:	23
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STEAMPENDERS both	

(D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
AGCCTGCTTT TTTGTACAAA GTTGG	25
(2) INFORMATION FOR SEQ ID NO:13:	
(1) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
AGCCTGCTT CTTGTACAAA GTTGG	2.5
	23
(2) INFORMATION FOR SEQ ID NO:14:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(x=' SEQUENCE DESCRIPTION: SEQ ID NO:14:	
ACCCAGCTTT CTTGTACAAA GTTGG	25
(2) INFORMATION FOR SEQ ID NO:15:	
(i' SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii' MOLECULE TYPE: cDNA	
(xi SEQUENCE DESCRIPTION: SEQ ID NO:15:	
GITCAGCITT TTTGTACAAA GTTGG	25

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(2)	NFORMATION FOR SEQ ID NO:16:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
	ii) MOLECULE TYPE: cDNA	
	xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
GTT	GCTTT CTTGTACAAA GTTGG	25
(2)	NFORMATION FOR SEQ ID NO:17:	
	(i) SEQUENCE CHARACTERISTICS: (A) LEMGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
	ii) MOLECULE TYPE: cDNA	
	xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
CCA	ACAAA CGCGTCCATG GAATTACACT TTAATTTAG	39
(2)	NFORMATION FOR SEQ ID NO:18:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
	ii) MOLECULE TYPE: cDNA	
	(i) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
CCA	ACAAG TCGACGCATG CCGACAGCCT TCCAAATGT	39
(2)	NFORMATION FOR SEQ ID NO:19:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 46 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
	1) MOLECULE TYPE: cDNA	

(X1) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
GGCCGATTAC GATATCCCAA CGACCGAAAA CCTGTATTTT CAGGGT	46
(2) INFORMATION FOR SEQ ID NO:20:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
CAGGTTTTCG GTCGTTGGGA TATCGTAATC	30
(2) INFORMATION FOR SEQ ID NO:21:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 47 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
GGCCAGATTA CGATATCCCA ACGACCGAAA ACCTGTATTT TCAGGGT	47
(2) INFORMATION FOR SEQ ID NO:22:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:	
CAGGITITICG GICGITIGGGA TATCGIAATC T	31
(2) INFORMATION FOR SEQ ID NO:23:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 48 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both	

(D) TOPOLOGY: both

(ii)	MOLECULE TYPE: cDNA	
	SEQUENCE DESCRIPTION: SEQ ID NO:23:	
	ATT ACGATATCCC AACGACCGAA AACCTGTATT TTCAGGGT	48
(2) INFO	RMATION FOR SEQ ID NO:24:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 32 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:24:	
CAGGTTTT	CG GTCGTTGGGA TATCGTAATC TT	32
(2) INFO	RMATION FOR SEQ ID NO:25:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 base pairs (B) TYFE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:25:	
ACCGTTT	ACG TGGAC	15
(2) INFO	DRMATION FOR SEQ ID NO:26:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:26:	
TCGAGTC	CAC GTAAACGGTT CCCACTTATT A	31

(2) INFORMATION FOR SEQ ID NO:27:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDMESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:	
UAUUUUCAGG GUATGGAGAA AAAAATCACT GGATATACC	39
(2) INFORMATION FOR SEQ ID NO:28:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:	
UCCCACUUAU UACGCCCCGC CCTGCCACTC ATC	33
(2) INFORMATION FOR SEQ ID NO:29:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:	
UAUUUUCAGG GUATGCCTGT TCTGGAAAAC CGG	33
(2) INFORMATION FOR SEQ ID NO:30:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:	
UCCCACUUAU UATTTCAGCC CCAGGGCGGC TTTC	34
(2) INFORMATION FOR SEQ ID NO:31:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:	
AGCCTGCTTT TTTATACTAA CTTGA	25
(2) INFORMATION FOR SEQ ID NO:32:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:	
TCAAGTTAGT ATAAAAAAGC AGGCT	25
(2) INFORMATION FOR SEQ ID NO:33:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDMESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:	
ACAAGTTTGT ACAAAAAAGC AGGCT	25
(2) INFORMATION FOR SEQ ID NO:34:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRENDEDUNGS, both	

(b) TOPOLOGI: Both	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:	
ACAAGTTTGT ACAAGAAAGC AGGCT	25
(2) INFORMATION FOR SEQ ID NO:35:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: both	
(ii) MOLECULE TYPE: cDNA	

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:

ACCACITTGT ACAAGAAAGC TGGGT